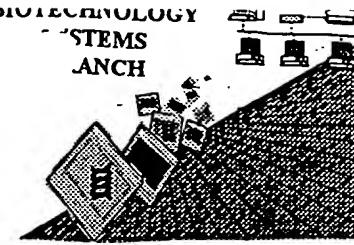


PCT09



CRF Problem Report

The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer readable form (CRF):

Application Serial Number: 09 | 937,204A
Filing Date: 09/21/2001
Date Processed by STIC: 11/25/02

STIC Contact: Mark Spencer, 703-308-4212

Nature of Problem:

The CRF (was):

Melted

- (circle one) Damaged or Unreadable (for Unreadable, see attached)
 Blank (no files on CRF) (see attached)
 Empty file (filename present, but no bytes in file) (see attached)
 Virus-infected. Virus name: _____ The STIC will not process the CRF.
 Not saved in ASCII text
 Sequence Listing was embedded in the file. According to Sequence Rules,
submitted file should only be the Sequence Listing.
 Did not contain a Sequence Listing. (see attached sample)
 Other:

**PLEASE USE THE CHECKER VERSION 3.1 PROGRAM TO REDUCE ERRORS.
SEE BELOW FOR ADDRESS:**

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service , or other delivery service to: U.S. Patent and Trademark Office,
Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



Does Not Comply
Corrected Diskette Needed

PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/937,204

DATE: 04/25/2002
TIME: 11:37:43

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt
Output Set: N:\CRF3\04252002\I937204.raw

```

4 <110> APPLICANT: Fritig, Bernard
5     Toquin, Valerie
6     Geoffroy, Pierrette
7     Legrand, Michel
8     Kauffmann, Serge
11 <120> TITLE OF INVENTION: INDUCIBLE COMTII PROMOTER, CHIMERA GENE
12     CONTAINING SAME AND TRANSFORMED PLANTS
15 <130> FILE REFERENCE: A34638-PCT-USA
17 <140> CURRENT APPLICATION NUMBER: US 09/937,204
18 <141> CURRENT FILING DATE: 2000-03-22
20 <150> PRIOR APPLICATION NUMBER: PCT/FR99/03700
21 <151> PRIOR FILING DATE: 1999-03-22
23 <150> PRIOR APPLICATION NUMBER: PCT/FR99/07646
24 <151> PRIOR FILING DATE: 1999-06-11
26 <160> NUMBER OF SEQ ID NOS: 26
28 <170> SOFTWARE: FastSEQ for Windows Version 4.0

```

ERRORED SEQUENCES

```

712 <210> SEQ ID NO: 26
713 <211> LENGTH: 30
714 <212> TYPE: DNA
715 <213> ORGANISM: Artificial Sequence
717 <220> FEATURE:
718 <223> OTHER INFORMATION: Synthetic Oligonucleotide PS14
720 <400> SEQUENCE: 26
    721 acgcgtcgac cagtggtag ttttagctgtc
E--> 734 ny02:358522.1 -delete

```

30

VARIABLE LOCATION SUMMARY
PATENT APPLICATION: US/09/937,204
DATE: 04/25/2002
TIME: 11:37:44

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt
Output Set: N:\CRF3\04252002\I937204.raw

Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which
residue n or Xaa represents.

Seq#:26; N Pos. 31

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/937,204

DATE: 04/25/2002
TIME: 11:37:44

Input Set : A:\#358522 v1 - A34638-PCT-USA Sequence Listing.txt
Output Set: N:\CRF3\04252002\I937204.raw

L:44 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:47 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:51 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:55 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:59 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:63 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:67 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:71 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:75 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:79 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:82 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:86 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:89 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:92 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:96 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:1
L:143 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:146 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:149 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:152 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:447 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:12
L:536 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:14
L:734 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:26
L:734 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:26
L:734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:30
L:734 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:34 SEQ:26
L:734 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10
L:734 M:112 C: (48) String data converted to lower case,
L:734 M:252 E: No. of Seq. differs, <211> LENGTH:Input:30 Found:34 SEQ:26